Ref #	Hits	Search Query	DBs	Default Operator	Plurals	Time Stamp
L1	2	"5981190".pn.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 09:44
L2	893	(karyotype or karyotyping and aneuploidy) and eukaryot\$	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 10:17
L3	587	L2 and @ad<="20021155"	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 10:21
L4	28	L3 and ((test and (reference or control)) adj cell)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 10:17
L5	25	L4 and ((genomic or tag or nucleotide or (nucleic adj acid)) adj sequence)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 10:18
L6	0	L5 and ((200kb or ("200" adj kb)) adj window)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 10:18
L7	0	L5 and ((200kb or ("200" adj kb)))	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 10:16
L8	0	L4 and ((200kb or ("200" adj kb)))	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 10:16
L9	85	L3 and ((200kb or ("200" adj kb)))	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 10:18

L10	0	L9 and ((test and (reference or control)) adj cell)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 10:16
L11	85	L9 and ((genomic or tag or nucleotide or (nucleic adj acid)) adj sequence)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 10:16
L12	1832	(karyotype or karyotyping or aneuploidy) and eukaryot\$	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 10:19
L13	1272	L12 and @ad<="20021155"	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 10:17
L14	117	L13 and ((test and (reference or control)) adj cell)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 10:25
L15	113	L14 and ((genomic or tag or nucleotide or (nucleic adj acid)) adj sequence)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 10:18
L16	0	L15 and ((200kb or ("200" adj kb)) adj window)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 10:18
L17	0	L15 and ((200kb or ("200" adj kb)))	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 10:19
L18	0	L14 and ((200kb or ("200" adj kb)))	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 10:21
L19	125	L13 and ((200kb or ("200" adj kb)))	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 10:19

						•
L20	266	(method same (karyotype or karyotyping)) and eukaryot\$	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 10:43
L21	147	L20 and @ad<="20021155"	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 10:21
L22	4	L21 and ((200kb or ("200" adj kb)))	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 10:44
L23	11	L20 and ((200kb or ("200" adj kb)))	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 10:24
L24	0	L22 and ((test and (reference or control)) adj cell)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 10:25
L25	2	L23 and ((test and (reference or control)) adj cell)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 10:25
L26	75	(method same (karyotype or karyotyping)same (genome or genomic)) and eukaryot\$	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON.	2006/05/01 10:49
L27	8	L26 and ((200kb or ("200" adj kb)))	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 10:45
L28	1	L26 and ((count or enumerate) with (tags or pieces or probes))	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 10:45
L29		L26 and ((count or enumerate or tabulate) with (tags or pieces or probes))	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 10:49

L30	0	L29 and ((200kb or ("200" adj kb)))	US-PGPUB;	OR	ON	2006/05/01 10:57
			USPAT; USOCR; EPO; JPO; DERWENT			2300,03,01 10.37
L31	765	(method same (karyotype or karyotyping))	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 10:49
L32	30	L31 and ((count or enumerate or tabulate) with (tags or pieces or probes))	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 11:06
L33	4	L32 and ((200kb or ("200" adj kb)))	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 10:49
L34	114	L31 and ((sequence adj tag) or (nucleotide adj tag) or GST or (genomic adj sequence adj tag))	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 11:02
L35	25	L34 and ((restriction adj endonuclease adj recognition adj site) or (restriction adj endonuclease adj recognition) or (restriction adj enzyme adj recognition adj site) or (restriction adj enzyme adj site))	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 11:04
L36	1	L35 and (((test adj cell) or (test adj genome)) and ((reference adj cell) or (reference adj cell)))	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 11:02
L39	0	L34 and ((((200kb or ("200" adj kb)) or (40kb or ("40" adj kb)) or (600kb or ("600" adj kb)) or (4Mb or ("4" adj Mb)))) adj (window or span))	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 11:09
L40	55	L31 and (((200kb or ("200" adj kb)) or (40kb or ("40" adj kb)) or (600kb or ("600" adj kb)) or (4Mb or ("4" adj Mb))))	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 11:01

L41	13	L40 and (((test adj cell) or (test adj genome)) and ((reference adj cell) or (reference adj cell)))	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 11:02
L42	12	L41 and (((sequence adj tag) or (nucleotide adj tag) or GST or (genomic adj sequence adj tag)) or ((genomic or tag or nucleotide or (nucleic adj acid)) adj sequence))	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 11:04
L44	101	L31 and ((restriction adj endonuclease adj recognition adj site) or (restriction adj endonuclease adj recognition) or (restriction adj enzyme adj recognition adj site) or (restriction adj enzyme adj site) or SacI or NlaIII or BcgI or MmeI)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 11:06
L45	1	L44 and ((count or enumerate or tabulate) with (tags or pieces or probes))	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 11:12
L46	0	L44 and ((((200kb or ("200" adj kb)) or (40kb or ("40" adj kb)) or (600kb or ("600" adj kb)) or (4Mb or ("4" adj Mb)))) adj (window or span))	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 11:10
L47	3	L44 and ((((200kb or ("200" adj kb)) or (40kb or ("40" adj kb)) or (600kb or ("600" adj kb)) or (4Mb or ("4" adj Mb)))))	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 11:10
L48	77	L44 and ((hereditary adj (disorder or disease)) or (infectious adj disease) or cancer)	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 11:11
L49	0	L48 and ((count or enumerate or tabulate) with (tags or pieces or probes))	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	ON	2006/05/01 11:12

Trying 31060000009999...Open DIALOG INFORMATION SERVICES PLEASE LOGON: ****** HHHHHHHH SSSSSSSS? ### Status: Signing onto Dialog ******* ENTER PASSWORD: ****** HHHHHHHH SSSSSSS? ****** ### Status: Login successfulWelcome to DIALOG Dialog level 05.11.05D Last logoff: 26apr06 13:52:54 Logon file405 01may06 10:17:16 *** ANNOUNCEMENTS *** NEW FILES RELEASED ***Regulatory Affairs Journals (File 183) ***Index Chemicus (File 302) ***Inspec (File 202) RESUMED UPDATING ***File 141, Reader's Guide Abstracts RELOADS COMPLETED ***File 516, D&B--Dun's Market Identifiers ***File 523, D&B European Dun's Market Identifiers ***File 531, American Business Directory *** MEDLINE has been reloaded with the 2006 MeSH (Files 154 & 155) *** The 2005 reload of the CLAIMS files (Files 340, 341, 942) is now available online. Chemical Structure Searching now available in Prous Science Drug Data Report (F452), Prous Science Drugs of the Future (F453), IMS R&D Focus (F445/955), Pharmaprojects (F128/928), Beilstein Facts (F390), Derwent Chemistry Resource (F355) and Index Chemicus (File 302). >>>For the latest news about Dialog products, services, content<<< >>>and events, please visit What's New from Dialog at <<< >>>http://www.dialog.com/whatsnew/. You can find news about<<< >>>a specific database by entering HELP NEWS <file number>.<< SYSTEM: HOME Cost is in DialUnits Menu System II: D2 version 1.7.9 term=ASCII *** DIALOG HOMEBASE(SM) Main Menu *** Information: Announcements (new files, reloads, etc.) 2. Database, Rates, & Command Descriptions 3. Help in Choosing Databases for Your Topic Customer Services (telephone assistance, training, seminars, etc.)
 Product Descriptions Connections: 6. DIALOG(R) Document Delivery 7. Data Star(R)

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Enter an option number to view information or to connect to an online service. Enter a BEGIN command plus a file number to search a database (e.g., B1 for ERIC).

Terminal set to DLINK

*** DIALOG HOMEBASE(SM) Main Menu ***

Information:

- 1. Announcements (new files, reloads, etc.)
- 2. Database, Rates, & Command Descriptions
- 3. Help in Choosing Databases for Your Topic
- 4. Customer Services (telephone assistance, training, seminars, etc.)
- 5. Product Descriptions

Connections:

- 6. DIALOG(R) Document Delivery
- 7. Data Star(R)
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/H = Help/L = Logoff/NOMENU = Command Mode

Enter an option number to view information or to connect to an online service. Enter a BEGIN command plus a file number to search a database (e.g., Bl for ERIC). ? b biosci

>>> 44 is unauthorized

76 is unauthorized >>>

>>>2 of the specified files are not available Olmay06 10:17:23 User276741 Session D133.1

\$0.00 0.214 DialUnits FileHomeBase

\$0.00 Estimated cost FileHomeBase \$0.02 TELNET \$0.02 Estimated cost this search

\$0.02 Estimated total session cost 0.214 DialUnits

SYSTEM:OS - DIALOG OneSearch

File 5:Biosis Previews(R) 1969-2006/Apr W4

(c) 2006 BIOSIS

File 24:CSA Life Sciences Abstracts 1966-2006/Mar

(c) 2006 CSA.

File 28:Oceanic Abstracts 1966-2006/Mar

(c) 2006 CSA.

File 34:SciSearch(R) Cited Ref Sci 1990-2006/Apr W3

(c) 2006 Inst for Sci Info

File 35: Dissertation Abs Online 1861-2006/Apr.

(c) 2006 ProQuest Info&Learning

40:Enviroline(R) 1975-2006/Mar File

File 41:Pollution Abstracts 1966-2006/Mar

(c) 2006 CSA.

File 50:CAB Abstracts 1972-2006/Mar

(c) 2006 CAB International

File 65:Inside Conferences 1993-2006/Apr 28

(c) 2006 BLDSC all rts. reserv.

File 71:ELSEVIER BIOBASE 1994-2006/Apr W5

```
(c) 2006 Elsevier Science B.V.
       73:EMBASE 1974-2006/May 01
         (c) 2006 Elsevier Science B.V.
       91:MANTIS(TM) 1880-2006/Feb
         2006 (c) Action Potential
       94:JICST-EPlus 1985-2006/Feb W1
         (c) 2006 Japan Science and Tech Corp(JST)
       98:General Sci Abs 1984-2004/Dec
         (c) 2005 The HW Wilson Co.
  File 110:WasteInfo 1974-2002/Jul
         (c) 2002 AEA Techn Env.
*File 110: This file is closed (no updates)
  File 135: NewsRx Weekly Reports 1995-2006/Apr W4
         (c) 2006 NewsRx
  File 136:BioEngineering Abstracts 1966-2006/Mar
         (c) 2006 CSA.
  File 143:Biol. & Agric. Index 1983-2006/Mar
         (c) 2006 The HW Wilson Co
  File 144: Pascal 1973-2006/Apr W2
         (c) 2006 INIST/CNRS
  File 155:MEDLINE(R) 1951-2006/May 03
         (c) format only 2006 Dialog
  File 164:Allied & Complementary Medicine 1984-2006/Apr
         (c) 2006 BLHCIS
  File 172:EMBASE Alert 2006/May 01
         (c) 2006 Elsevier Science B.V.
  File 185: Zoological Record Online (R) 1978-2006/May
         (c) 2006 BIOSIS
  File 357: Derwent Biotech Res. 1982-2006/Apr W4
         (c) 2006 Thomson Derwent & ISI
  File 369: New Scientist 1994-2006/Sep W1
         (c) 2006 Reed Business Information Ltd.
  File 370:Science 1996-1999/Jul W3
         (c) 1999 AAAS
*File 370: This file is closed (no updates). Use File 47 for more current
information.
  File 391:Beilstein Reactions 2006/Q1
         (c) 2005 Beilstein GmbH
  File 434:SciSearch(R) Cited Ref Sci 1974-1989/Dec
         (c) 1998 Inst for Sci Info
  File 467:ExtraMED(tm) 2000/Dec
         (c) 2001 Informania Ltd.
*File 467: F467 will close on February 1, 2006.
                                                                        7.
     Set Items Description
          _____
? s (method (s) (karyotype or karyotyping))
         8990351 METHOD
         131204 KARYOTYPE
           56255 KARYOTYPING
           5934 (METHOD (S) (KARYOTYPE OR KARYOTYPING))
      S1
? s s1 and ((test and (reference or control)) (w) cell)
Processing
Processed 10 of 29 files ...
Processing
Processed 20 of 29 files ...
Completed processing all files
           5934 S1
         4851845 TEST
         1161336 REFERENCE
         9587432 CONTROL
```

```
13879396 CELL
           2000 (TEST AND (REFERENCE OR CONTROL))(W)CELL
      S2
              3 S1 AND ((TEST AND (REFERENCE OR CONTROL)) (W) CELL)
? rd
>>>Duplicate detection is not supported for File 391.
>>>Records from unsupported files will be retained in the RD set.
      s3
              3 RD (unique items)
? s s3 and ((genomic or tag or nucleotide or (nucleic (w) acid)) (w) sequence)
Processing
Processed 10 of 29 files ...
Processing
Completed processing all files
              3 S3
         579895 GENOMIC
          90667 TAG
        1331728 NUCLEOTIDE
         661864 NUCLEIC
       11986172 ACID
         541592 NUCLEIC(W)ACID
         3492911 SEQUENCE
         655734
                 (((GENOMIC OR TAG) OR NUCLEOTIDE) OR
                 NUCLEIC (W) ACID) (W) SEQUENCE
      S4
               1 S3 AND ((GENOMIC OR TAG OR NUCLEOTIDE OR (NUCLEIC (W)
                 ACID)) (W) SEQUENCE)
? t s4/free
 4/8/1
           (Item 1 from file: 357)
0343662 DBR Accession No.: 2004-15954
Digital karyotyping a genome of a test eukaryotic cell comprises isolating
   and enumerating short sequence tags from specific genomic loci and
    comparing the sequence tags to a genome of a reference cell - using
   bioinformatics for human cancer cell karyotyping for use in disease
   diagnosis, therapy and genomics 2004
? s sl and ((genomic or tag or nucleotide or (nucleic (w) acid)) (w) sequence)
Processing
Processed 20 of 29 files ...
Completed processing all files
            5934 S1
         579895 GENOMIC
           90667 TAG
        1331728 NUCLEOTIDE
         661864 NUCLEIC
       11986172 ACID
         541592 NUCLEIC (W) ACID
        3492911 SEQUENCE
         655734 (((GENOMIC OR TAG) OR NUCLEOTIDE) OR
                 NUCLEIC (W) ACID) (W) SEQUENCE
      S5
              74 S1 AND ((GENOMIC OR TAG OR NUCLEOTIDE OR (NUCLEIC (W)
                 ACID)) (W) SEQUENCE)
? s s5 and ((((200kb or ("200" (w) kb)) or (40kb or ("40" (w) kb)) or (600kb
or ("600" (w) kb)) or (4Mb or ("4" (w) Mb))) ))
Processing
Processed 10 of 29 files ...
Processing
Processed 20 of 29 files ...
Completed processing all files
             74 S5
             102 200KB
        1099242 200
```

```
343680 KB
            3102 200(W)KB
             108 40KB
         2802791
                 40
          343680 KB
            3626 40(W)KB
                 600KB
              31
          339800
                 600
          343680 KB
            1094 600(W)KB
             424
                 4MB
        20170163
          107987
                 MB
            2767
                 4 (W) MB
               1 S5 AND ((((200KB OR ("200" (W) KB)) OR (40KB OR ("40" (W)
      S 6
                  KB)) OR (600KB OR ("600" (W) KB)) OR (4MB OR ("4" (W)
                  MB)))))))
? t s6/free
 6/8/1
           (Item 1 from file: 357)
0343662
         DBR Accession No.: 2004-15954
Digital karyotyping a genome of a test eukaryotic cell comprises isolating
    and enumerating short sequence tags from specific genomic loci and
    comparing the sequence tags to a genome of a reference cell - using
    bioinformatics for human cancer cell karyotyping for use in disease
    diagnosis, therapy and genomics 2004
? s sl and ((count or enumerate or tabulate) (w)(tags or pieces or probes))
            5934 S1
          587149 COUNT
            8331 ENUMERATE
            1973 TABULATE
           49899 TAGS
          103655 PIECES
          455040 PROBES
                 ((COUNT OR ENUMERATE) OR TABULATE) (W) ((TAGS OR PIECES) OR
      S7
                 S1 AND ((COUNT OR ENUMERATE OR TABULATE) (W) (TAGS OR
                  PIECES OR PROBES))
? s sl and (((sequence (w) tag) or (nucleotide (w)tag) or GST or (genomic (w)
sequence (w) tag)) or ((genomic or tag or nucleotide or (nucleic (w) acid))
(w) sequence))
Processing
Processed 10 of 29 files ...
Completed processing all files
            5934 S1
         3492911 SEQUENCE
           90667 TAG
           18540 SEQUENCE (W) TAG
         1331728 NUCLEOTIDE
           90667 TAG
              53 NUCLEOTIDE (W) TAG
           61336 GST
          579895 GENOMIC
         3492911 SEQUENCE
           90667 TAG
               5 GENOMIC (W) SEQUENCE (W) TAG
          579895 GENOMIC
           90667
                 TAG
         1331728 NUCLEOTIDE
          661864 NUCLEIC
        11986172 ACID
```

```
541592 NUCLEIC (W) ACID
         3492911 SEQUENCE
          655734
                  (((GENOMIC OR TAG) OR NUCLEOTIDE) OR
                  NUCLEIC (W) ACID) (W) SEQUENCE
      S8
              75
                  S1 AND (((SEQUENCE (W) TAG) OR (NUCLEOTIDE (W) TAG) OR GST
                  OR (GENOMIC (W) SEQUENCE (W) TAG)) OR ((GENOMIC OR TAG OR
                  NUCLEOTIDE OR (NUCLEIC (W) ACID)) (W) SEQUENCE))
? s s8 and ((restriction (w) endonuclease (w) recognition (w) site) or
(restriction (w) endonuclease (w) recognition) or (restriction (w) enzyme (w)
recognition (w) site) or (restriction (w) enzyme (w) site) or SacI or NlaIII
or BcgI or MmeI)
Processing
Processed 10 of 29 files ...
Completed processing all files
              75 S8
          690035 RESTRICTION
          120352 ENDONUCLEASE
         1081484 RECOGNITION
         3171822 SITE
             257
                 RESTRICTION (W) ENDONUCLEASE (W) RECOGNITION (W) SITE
          690035 RESTRICTION
          120352 ENDONUCLEASE
         1081484 RECOGNITION
             664
                 RESTRICTION (W) ENDONUCLEASE (W) RECOGNITION
          690035 RESTRICTION
         4217353 ENZYME
         1081484 RECOGNITION
         3171822 SITE
             246 RESTRICTION (W) ENZYME (W) RECOGNITION (W) SITE
          690035 RESTRICTION
         4217353 ENZYME
         3171822 SITE
            1462 RESTRICTION(W) ENZYME(W) SITE
            3342 SACI
             582 NLAIII
             133
                 BCGI
              65
                  MMEI
      S 9
                  S8 AND ((RESTRICTION (W) ENDONUCLEASE (W) RECOGNITION (W)
                  SITE) OR (RESTRICTION (W) ENDONUCLEASE (W) RECOGNITION)
                  OR (RESTRICTION (W) ENZYME (W) RECOGNITION (W) SITE) OR
                  (RESTRICTION (W) ENZYME (W) SITE) OR SACI OR NLAIII OR
                  BCGI OR MMEI)
? ds
Set
        Items
                Description
S1
         5934
                (METHOD (S) (KARYOTYPE OR KARYOTYPING))
S2
                S1 AND ((TEST AND (REFERENCE OR CONTROL)) (W) CELL)
S3
                   (unique items)
            3
S4
                S3 AND ((GENOMIC OR TAG OR NUCLEOTIDE OR (NUCLEIC (W) ACID-
             )) (W) SEQUENCE)
S5
                S1 AND ((GENOMIC OR TAG OR NUCLEOTIDE OR (NUCLEIC (W) ACID-
             )) (W) SEQUENCE)
                S5 AND ((((200KB OR ("200" (W) KB)) OR (40KB OR ("40" (W) -
S 6
            KB)) OR (600KB OR ("600" (W) KB)) OR (4MB OR ("4" (W) MB))) )
S7
                S1 AND ((COUNT OR ENUMERATE OR TABULATE) (W)(TAGS OR PIECES
             OR PROBES))
S8
               S1 AND (((SEQUENCE (W) TAG) OR (NUCLEOTIDE (W)TAG) OR GST -
             OR (GENOMIC (W) SEQUENCE (W) TAG)) OR ((GENOMIC OR TAG OR NUC-
             LEOTIDE OR (NUCLEIC (W) ACID)) (W) SEQUENCE))
S9
               S8 AND ((RESTRICTION (W) ENDONUCLEASE (W) RECOGNITION (W) -
```

```
SITE) OR (RESTRICTION (W) ENDONUCLEASE (W) RECOGNITION) OR (R-
             ESTRICTION (W) ENZYME (W) RECOGNITION (W) SITE) OR (RESTRICTI-
             ON (W) ENZYME (W) SITE) OR SACI OR NLAIII OR BCGI OR MMEI)
>>>Duplicate detection is not supported for File 391.
>>>Records from unsupported files will be retained in the RD set.
               2 RD (unique items)
? t s10/medium/all
            (Item 1 from file: 357)
DIALOG(R) File 357: Derwent Biotech Res.
(c) 2006 Thomson Derwent & ISI. All rts. reserv.
0379012 DBR Accession No.: 2005-24718
                                           PATENT
Restriction enzyme tag concatemer nucleic acid library useful for
    identifying pathogen and detecting expression of desired transcripts,
    in biological sample, comprises several concatemers having Type IIB
    restriction enzyme tags - restriction enzyme tag concatemer DNA library
    for use in disease diagnosis and genome mapping
AUTHOR: MEYERSON M L; TENGS T
PATENT ASSIGNEE: DANA FARBER CANCER INST 2005
PATENT NUMBER: WO 200579357 PATENT DATE: 20050901 WPI ACCESSION NO.:
    2005-591909 (200560)
PRIORITY APPLIC. NO.: US 545047 APPLIC. DATE: 20040217
NATIONAL APPLIC. NO.: WO 2005US4571 APPLIC. DATE: 20050215
LANGUAGE: English
            (Item 2 from file: 357)
DIALOG(R) File 357: Derwent Biotech Res.
(c) 2006 Thomson Derwent & ISI. All rts. reserv.
0343662 DBR Accession No.: 2004-15954
                                           PATENT
Digital karyotyping a genome of a test eukaryotic cell comprises isolating
    and enumerating short sequence tags from specific genomic loci and
    comparing the sequence tags to a genome of a reference cell - using
    bioinformatics for human cancer cell karyotyping for use in disease
    diagnosis, therapy and genomics
AUTHOR: WANG T; VELCULESCU V; KINZLER K; VOGELSTEIN B ·
PATENT ASSIGNEE: UNIV JOHNS HOPKINS 2004
PATENT NUMBER: US 20040096892 PATENT DATE: 20040520 WPI ACCESSION NO.:
2004-389156 (200436)
PRIORITY APPLIC. NO.: US 705874 APPLIC. DATE: 20031113
NATIONAL APPLIC. NO.: US 705874 APPLIC. DATE: 20031113
LANGUAGE: English
? s s8 and ((hereditary (w) (disorder or disease)) or (infectious (w) disease)
Processed 10 of 29 files ...
Completed processing all files
         75 S8
247964 HEREDITARY
1979919 DISORDER
        12746172 DISEASE
27057 HEREDITARY(W)(DISORDER OR DISEASE)
1157491 INFECTIOUS
```

? rd

S10

10/3/1

10/3/2

or cancer) Processing

Processing

12746172 DISEASE

```
299220 INFECTIOUS (W) DISEASE
         4037448 CANCER
     S11
             19 S8 AND ((HEREDITARY (W) (DISORDER OR DISEASE)) OR
                  (INFECTIOUS (W) DISEASE) OR CANCER)
? rd
>>>Duplicate detection is not supported for File 391.
>>>Records from unsupported files will be retained in the RD set.
     S12
              17 RD (unique items)
? s s12 and ((count or enumerate or tabulate) (w)(tags or pieces or probes))
              17 S12
          587149 COUNT
            8331 ENUMERATE
            1973 TABULATE
           49899 TAGS
          103655 PIECES
          455040 PROBES
                 ((COUNT OR ENUMERATE) OR TABULATE) (W) ((TAGS OR PIECES) OR
                  PROBES)
     S13
               O S12 AND ((COUNT OR ENUMERATE OR TABULATE) (W) (TAGS OR
                  PIECES OR PROBES))
? ds
Set
        Items
                Description
         5934
S1
                (METHOD (S) (KARYOTYPE OR KARYOTYPING))
S2
                S1 AND ((TEST AND (REFERENCE OR CONTROL)) (W) CELL)
                RD (unique items)
S3
            3
                S3 AND ((GENOMIC OR TAG OR NUCLEOTIDE OR (NUCLEIC (W) ACID-
S 4
            )) (W) SEQUENCE)
           74
                S1 AND ((GENOMIC OR TAG OR NUCLEOTIDE OR (NUCLEIC (W) ACID-
S 5
            )) (W) SEQUENCE)
               S5 AND ((((200KB OR ("200" (W) KB)) OR (40KB OR ("40" (W) -
S6
             KB)) OR (600KB OR ("600" (W) KB)) OR (4MB OR ("4" (W) MB))) )
                S1 AND ((COUNT OR ENUMERATE OR TABULATE) (W) (TAGS OR PIECES
S7
              OR PROBES))
               S1 AND (((SEQUENCE (W) TAG) OR (NUCLEOTIDE (W) TAG) OR GST -
S8
             OR (GENOMIC (W) SEQUENCE (W) TAG)) OR ((GENOMIC OR TAG OR NUC-
             LEOTIDE OR (NUCLEIC (W) ACID)) (W) SEQUENCE))
                S8 AND ((RESTRICTION (W) ENDONUCLEASE (W) RECOGNITION (W) -
S9
             SITE) OR (RESTRICTION (W) ENDONUCLEASE (W) RECOGNITION) OR (R-
             ESTRICTION (W) ENZYME (W) RECOGNITION (W) SITE) OR (RESTRICTI-
             ON (W) ENZYME (W) SITE) OR SACI OR NLAIII OR BCGI OR MMEI)
S10
                   (unique items)
                RD
                S8 AND ((HEREDITARY (W) (DISORDER OR DISEASE)) OR (INFECTI-
S11
             OUS (W) DISEASE) OR CANCER)
$12
                RD (unique items)
S13
                S12 AND ((COUNT OR ENUMERATE OR TABULATE) (W) (TAGS OR PIEC-
             ES OR PROBES))
? s (dimers or ditags) and (eukaryot?? (s) genome)
          128715 DIMERS
              78 DITAGS
          593355 EUKARYOT??
          793687
                 GENOME
           25771 EUKARYOT??(S)GENOME
             154 (DIMERS OR DITAGS) AND (EUKARYOT?? (S) GENOME)
? s s14 and ((restriction (w) endonuclease (w) recognition (w) site) or
(restriction (w) endonuclease (w) recognition) or (restriction (w) enzyme (w)
recognition (w) site) or (restriction (w) enzyme (w) site) or SacI or NlaIII
or BcgI or MmeI)
```

```
Processing
Processed 20 of 29 files ...
Completed processing all files
             154 S14
          690035 RESTRICTION
          120352 ENDONUCLEASE
         1081484 RECOGNITION
         3171822 SITE
             257 RESTRICTION (W) ENDONUCLEASE (W) RECOGNITION (W) SITE
          690035 RESTRICTION
          120352 ENDONUCLEASE
         1081484 RECOGNITION
             664 RESTRICTION (W) ENDONUCLEASE (W) RECOGNITION
          690035 RESTRICTION
         4217353 ENZYME
         1081484 RECOGNITION
         3171822 SITE
             246 RESTRICTION(W) ENZYME(W) RECOGNITION(W) SITE
          690035 RESTRICTION
         4217353 ENZYME
                 SITE
         3171822
            1462 RESTRICTION (W) ENZYME (W) SITE.
            3342
                 SACI
             582 NLAIII
             133
                 BCGI
              65
                 MMEI
     S15
                 S14 AND ((RESTRICTION (W) ENDONUCLEASE (W) RECOGNITION
                  (W) SITE) OR (RESTRICTION (W) ENDONUCLEASE (W)
                  RECOGNITION) OR (RESTRICTION (W) ENZYME (W) RECOGNITION
                  (W) SITE) OR (RESTRICTION (W) ENZYME (W) SITE) OR SACI OR
                  NLAIII OR BCGI OR MMEI)
? rd
>>>Duplicate detection is not supported for File 391.
>>>Records from unsupported files will be retained in the RD set.
             2 RD (unique items)
? t s16/medium/all
           (Item 1 from file: 98)
DIALOG(R) File 98: General Sci Abs
(c) 2005 The HW Wilson Co. All rts. reserv.
04050517
           H.W. WILSON RECORD NUMBER: BGSA99050517 (USE FORMAT 7 FOR
FULLTEXT)
Giant viruses infecting algae.
AUGMENTED TITLE: review
Van Etten, James L
Meints, Russel H
Annual Review of Microbiology v. 53 (1999) p. 447-94
SPECIAL FEATURES: bibl il ISSN: 0066-4227
LANGUAGE: English
COUNTRY OF PUBLICATION: United States
WORD COUNT: 22140
          (Item 1 from file: 357)
16/3/2
DIALOG(R) File 357: Derwent Biotech Res.
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```

0343662 DBR Accession No.: 2004-15954

```
Digital karyotyping a genome of a test eukaryotic cell comprises
    isolating and enumerating short sequence tags from specific genomic
    loci and comparing the sequence tags to a genome of a reference cell
    - using bioinformatics for human cancer cell karyotyping for use in
    disease diagnosis, therapy and genomics
AUTHOR: WANG T; VELCULESCU V; KINZLER K; VOGELSTEIN B PATENT ASSIGNEE: UNIV JOHNS HOPKINS 2004
PATENT NUMBER: US 20040096892 PATENT DATE: 20040520 WPI ACCESSION NO.:
2004-389156 (200436)
PRIORITY APPLIC. NO.: US 705874 APPLIC. DATE: 20031113
NATIONAL APPLIC. NO.: US 705874 APPLIC. DATE: 20031113
LANGUAGE: English
? s (concatamer and (dimers or ditags))
             297 CONCATAMER
          128715 DIMERS
              78 DITAGS
              12 (CONCATAMER AND (DIMERS OR DITAGS))
? s s17 and ((restriction (w) endonuclease (w) recognition (w) site) or
(restriction (w) endonuclease (w) recognition) or (restriction (w) enzyme (w)
recognition (w) site) or (restriction (w) enzyme (w) site) or SacI or NlaIII
or BcgI or MmeI)
Processing
Processed 20 of 29 files ...
Completed processing all files
              12 S17
          690035 RESTRICTION
          120352 ENDONUCLEASE
         1081484 RECOGNITION
         3171822 SITE
             257 RESTRICTION (W) ENDONUCLEASE (W) RECOGNITION (W) SITE
          690035 RESTRICTION
          120352 ENDONUCLEASE
         1081484 RECOGNITION
             664 RESTRICTION (W) ENDONUCLEASE (W) RECOGNITION
          690035 RESTRICTION
         4217353 ENZYME
         1081484 RECOGNITION
         3171822 SITE
             246 RESTRICTION (W) ENZYME (W) RECOGNITION (W) SITE
          690035 RESTRICTION
         4217353 ENZYME
         3171822 SITE
            1462 RESTRICTION (W) ENZYME (W) SITE
            3342 SACI
             582 NLAIII
             133 BCGI
              65 MMEI
     S18
               3 S17 AND ((RESTRICTION (W) ENDONUCLEASE (W) RECOGNITION
                  (W) SITE) OR (RESTRICTION (W) ENDONUCLEASE (W)
                  RECOGNITION) OR (RESTRICTION (W) ENZYME (W) RECOGNITION
                  (W) SITE) OR (RESTRICTION (W) ENZYME (W) SITE) OR SACI OR
                  NLAIII OR BCGI OR MMEI)
? rd
>>>Duplicate detection is not supported for File 391.
>>>Records from unsupported files will be retained in the RD set.
     S19
              3 RD (unique items)
? t s19/medium, k/all
 19/K/1
            (Item 1 from file: 357)
```

DIALOG(R) File 357: Derwent Biotech Res. (c) 2006 Thomson Derwent & ISI. All rts. reserv.

0367226 DBR Accession No.: 2005-12933 PATENT

New isolated oligonucleotides comprising at least one ditag, useful for gene identification signature analysis or for gene discovery or genome mapping - oligonucleotide isolation for use in gene mapping and gene discovery

AUTHOR: RUAN Y; NG P; WEI C

PATENT ASSIGNEE: AGENCY SCI TECHNOLOGY and RES 2005

PATENT NUMBER: US 20050059022 PATENT DATE: 20050317 WPI ACCESSION NO.:

2005-240897 (200525)

PRIORITY APPLIC. NO.: US 664234 APPLIC. DATE: 20030917 NATIONAL APPLIC. NO.: US 664234 APPLIC. DATE: 20030917

LANGUAGE: English

- ...ABSTRACT: It is selected from AarI, AceIII, AloI, BaeI, Bbr7I, BbvI, BbvII, BccI, Bce83I, BceAI, BcefI, BcgI, BciVI, BfiI, BinI, BplI, MmeI, etc. The ditag size is 12-60 bp. The ditag comprises 34-38 nucleotides and the size of each tag is determined by the use of restriction enzyme MmeI. The first and second tags have the same or a different number of nucleotides. The oligonucleotide consists of 1-1000 ditags. Preferred Vector: The vector further comprises the above-mentioned 2 adapter sequences flanking the ditag...
- ... including at least one ditag flanked by the 2 adapters. The method further comprises creating concatamer of ditags. The ditags are 1-1000 ditags. The method further comprises including the oligonucleotide comprising the ditag flanked by the adapters in...
- ... at least one oligonucleotide cited above, where splicing includes cleaving the full-length cDNA with **MmeI** which forms 3' overhanging tag ends, and ligating the two 5' and 3' termini tags...

19/K/2 (Item 2 from file: 357)

DIALOG(R) File 357: Derwent Biotech Res.

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0344040 DBR Accession No.: 2004-16332 PATENT

Obtaining a nucleotide sequence tags from termini of nucleic acids, useful for novel gene discoveries, comprises linking a first nucleic acid sequence to a linker sequence and cleaving it with a nucleic acid cleavage enzyme - gene expression profiling technique

AUTHOR: RUAN Y; WEI C

PATENT ASSIGNEE: AGENCY SCI TECHNOLOGY and RES 2004

PATENT NUMBER: WO 200450918 PATENT DATE: 20040617 WPI ACCESSION NO.:

2004-450747 (200442)

PRIORITY APPLIC. NO.: GB 200228289 APPLIC. DATE: 20021204 NATIONAL APPLIC. NO.: WO 2003SG255 APPLIC. DATE: 20031204

LANGUAGE: English

...ABSTRACT: site or the second recognition site, or both, comprises a Type IIS restriction enzyme recognition site. The first restriction enzyme comprises Mmel, or in which the first recognition site comprises a...

19/K/3 (Item 3 from file: 357)

DIALOG(R) File 357: Derwent Biotech Res.

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0343662 DBR Accession No.: 2004-15954 PATENT

Digital karyotyping a genome of a test eukaryotic cell comprises isolating and enumerating short sequence tags from specific genomic loci and comparing the sequence tags to a genome of a reference cell - using bioinformatics for human cancer cell karyotyping for use in disease diagnosis, therapy and genomics

AUTHOR: WANG T; VELCULESCU V; KINZLER K; VOGELSTEIN B

PATENT ASSIGNEE: UNIV JOHNS HOPKINS 2004

PATENT NUMBER: US 20040096892 PATENT DATE: 20040520 WPI ACCESSION NO.:

2004-389156 (200436)

PRIORITY APPLIC. NO.: US 705874 APPLIC. DATE: 20031113 NATIONAL APPLIC. NO.: US 705874 APPLIC. DATE: 20031113

LANGUAGE: English

- ...ABSTRACT: the genome of the test eukaryotic cell, the portions being defined by 1 or 2 restriction endonuclease recognition sites; (b) enumerating the sequence tags in the population to determine the number of individual...
- ...of a genome of a eukaryotic cell, the portions being defined by 1 or 2 restriction endonuclease recognition sites, where each of the sequence tags consists of a fixed number of nucleotides of...
- ... of the genome, the fixed number of nucleotides extending from at least one of the restriction endonuclease recognition sites; and (2) a concatamer of dimers cited above. BIOTECHNOLOGY Preferred Method: In karyotyping a genome of a test eukaryotic cell, the...
- ... endonuclease cleavage site at a second end of each portion. The first restriction endonuclease is **SacI** and the second restriction endonuclease is NlaIII. Recognition or cleavage by the first restriction endonuclease is sensitive to DNA methylation. The portions are defined by presence of a **BcgI** restriction endonuclease recognition site which is flanked by 12 nucleotides on either end. The method further comprises identifying aneuploidy...
- ... to cleave within the defined portions of the genome. The Type IIS restriction endonuclease is **MmeI**. The fixed number of nucleotides is 20-22. USE The method is useful for determining...
- ... manufacturers' protocols. For each sample 1microg of genomic DNA was sequentially digested with mapping enzyme SacI , ligated to 20-40 ng of biotinylated linker (5'-biotin-TTTGCAGAGGTTCGTAATC- GAGTTGGGTGAGCT(S EO ID NO...
- ... Coralville, Iowa) using T4 ligase (Invitrogen, Carlsbad, Calif.), and then digested with the fragmenting enzyme NlaIII. DNA fragments containing biotinylated linkers were isolated by binding to streptavidin coated magnetic beads (Dynal...
- ... steps were similar to those described for LongSAGE of cDNA (18). In brief, linkers containing MmeI recognition sites were ligated to captured DNA fragments, and tags were released with MmeI (University of Gdansk Center for Technology Transfer, Gdansk, Poland and New England Biolabs, Beverly, Mass.). The tags were ligated to form ditags, and the ditags were isolated and then ligated to form concatamers which were cloned into pZero (Invitrogen, Carlsbad, Calif.). Sequencing of concatamer clones was performed using the Big Dye Terminator v3.0 Kit (Applied Biosystems) and analyzed...

```
identifies the fragmenting enzyme site between ditags, extracts
    intervening tags, and records them in a database. Detailed protocols
    for performing Digital Karyotyping...
DESCRIPTORS: human colorectal cancer cell digital karyotyping, short
    sequence tag enumeration, restriction endonuclease SacI , N1aIII, DNA
   methylation, DNA copy number det., computer bioinformatic software,
    appl. genomics, cancer diagnosis, therapy...
? ds
Set
        Items
                Description
S1
         5934
                (METHOD (S) (KARYOTYPE OR KARYOTYPING))
S2
                S1 AND ((TEST AND (REFERENCE OR CONTROL)) (W) CELL)
                   (unique items)
S3
            3
S4
                S3 AND ((GENOMIC OR TAG OR NUCLEOTIDE OR (NUCLEIC (W) ACID-
             )) (W) SEQUENCE)
S5.
                S1 AND ((GENOMIC OR TAG OR NUCLEOTIDE OR (NUCLEIC (W) ACID-
             )) (W) SEQUENCE)
                S5 AND ((((200KB OR ("200" (W) KB)) OR (40KB OR ("40" (W) -
S6
             KB)) OR (600KB OR ("600" (W) KB)) OR (4MB OR ("4" (W) MB))) )
S7
                S1 AND ((COUNT OR ENUMERATE OR TABULATE) (W) (TAGS OR PIECES
              OR PROBES))
S8
                S1 AND (((SEQUENCE (W) TAG) OR (NUCLEOTIDE (W) TAG) OR GST -
             OR (GENOMIC (W) SEQUENCE (W) TAG)) OR ((GENOMIC OR TAG OR NUC-
             LEOTIDE OR (NUCLEIC (W) ACID)) (W) SEQUENCE))
S9
                S8 AND ((RESTRICTION (W) ENDONUCLEASE (W) RECOGNITION (W) -
             SITE) OR (RESTRICTION (W) ENDONUCLEASE (W) RECOGNITION) OR (R-
             ESTRICTION (W) ENZYME (W) RECOGNITION (W) SITE) OR (RESTRICTI-
             ON (W) ENZYME (W) SITE) OR SACI OR NLAIII OR BCGI OR MMEI)
S10
                   (unique items)
S11
                S8 AND ((HEREDITARY (W) (DISORDER OR DISEASE)) OR (INFECTI-
             OUS (W) DISEASE) OR CANCER)
S12
           17
                RD (unique items)
S13
                S12 AND ((COUNT OR ENUMERATE OR TABULATE) (W)(TAGS OR PIEC-
             ES OR PROBES))
S14
          154
                (DIMERS OR DITAGS) AND (EUKARYOT?? (S) GENOME)
S15
                S14 AND ((RESTRICTION (W) ENDONUCLEASE (W) RECOGNITION (W)
             SITE) OR (RESTRICTION (W) ENDONUCLEASE (W) RECOGNITION) OR (R-
             ESTRICTION (W) ENZYME (W) RECOGNITION (W) SITE) OR (RESTRICTI-
             ON (W) ENZYME (W) SITE) OR SACI OR NLAIII OR BCGI OR MMEI)
S16
            2
                   (unique items)
S17
           12
                (CONCATAMER AND (DIMERS OR DITAGS))
                S17 AND ((RESTRICTION (W) ENDONUCLEASE (W) RECOGNITION (W)
S18
             SITE) OR (RESTRICTION (W) ENDONUCLEASE (W) RECOGNITION) OR (R-
             ESTRICTION (W) ENZYME (W) RECOGNITION (W) SITE) OR (RESTRICTI-
             ON (W) ENZYME (W) SITE) OR SACI OR NLAIII OR BCGI OR MMEI)
S19
               RD (unique items)
? save temp
Temp SearchSave "TD231378228" stored
? logoff
       Olmay06 10:36:34 User276741 Session D133.2
           $22.12
                     3.749 DialUnits File5
           Estimated cost File5
    $22.12
            $5.16
                    0.832 DialUnits File24
           Estimated cost File24
    $5.16
            $0.86
                    0.139 DialUnits File28
    $0.86 Estimated cost File28
           $50.70
                    2.160 DialUnits File34
    $50.70 Estimated cost File34
```

\$1.41

0.344 DialUnits File35

... tags were extracted using the SAGE2000 software package, which

\$1.41 Estimated cost File35 0.117 DialUnits File40 \$0.84 \$0.84 Estimated cost File40 0.115 DialUnits File41 \$0.72 \$0.72 Estimated cost File41 \$3.98 0.864 DialUnits File50 \$3.98 Estimated cost File50 \$0.74 0.197 DialUnits File65 \$0.74 Estimated cost File65 1.091 DialUnits File71 \$9.60 \$9.60 Estimated cost File71 \$33.83 3.021 DialUnits File73 \$33.83 Estimated cost File73 \$0.49 0.113 DialUnits File91 \$0.49 Estimated cost File91 \$5.07 1.448 DialUnits File94 \$5.07 Estimated cost File94 \$1.11 0.261 DialUnits File98 \$1.45 1 Type(s) in Format 3 \$1.45 1 Types \$2.56 Estimated cost File98 \$0.53 0.092 DialUnits File110 \$0.53 Estimated cost File110 \$1.17 0.217 DialUnits File135 \$1.17 Estimated cost File135 \$0.96 0.155 DialUnits File136 \$0.96 Estimated cost File136 \$0.42 Estimated cost File143 \$10.58 2.351 DialUnits File144 \$10.58 Estimated cost File144 \$10.36 3.046 DialUnits File155 \$10.36 Estimated cost File155 \$0.33 0.096 DialUnits File164 \$0.33 Estimated cost File164 \$1.36 Estimated cost File172 0.213 DialUnits File185 \$1.31 \$1.31 Estimated cost File185 0.880 DialUnits File357 \$19.63 \$15.60 6 Type(s) in Format 3 \$0.00 2 Type(s) in Format 6 \$15.60 8 Types \$35.23 Estimated cost File357 \$0.34 0.098 DialUnits File369 \$0.34 Estimated cost File369 \$0.43 0.121 DialUnits File370 \$0.43 Estimated cost File370 \$0.00 1.223 DialUnits File391 \$0.00 Estimated cost File391 0.404 DialUnits File434 \$9.49 \$9.49 Estimated cost File434 0.106 DialUnits File467 \$0.68 \$0.68 Estimated cost File467 OneSearch, 29 files, 23.716 DialUnits FileOS \$5.33 TELNET \$216.60 Estimated cost this search \$216.62 Estimated total session cost 23.930 DialUnits